

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

101702,319
1540 -
11/17/03~

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual cPAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry directly to (EFFECTIVE 12/01/03):
 U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- 4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 4B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03



IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/702,319

DATE: 11/17/2003

TIME: 09:26:36

Input Set : A:\1036C5SEQLIST.txt

Output Set: N:\CRF4\11172003\J702319.raw

4 <110> APPLICANT: Perera, Ranjan

Rice, Stephen

Eagleton, Clare

9 <120> TITLE OF INVENTION: Compositions and Methods for the

10 Modification of Gene Expression

13 <130> FILE REFERENCE: 11000.1036c5

C--> 15 <140> CURRENT APPLICATION NUMBER: US/10/702,319

C--> 15 <141> CURRENT FILING DATE: 2003-11-06

15 <150> PRIOR APPLICATION NUMBER: U.S. No. 10/291,447

16 <151> PRIOR FILING DATE: 2002-11-08

18 <150> PRIOR APPLICATION NUMBER: U.S. No. 60/425,087

19 <151> PRIOR FILING DATE: 2002-11-08

21 <150> PRIOR APPLICATION NUMBER: U.S. No. 10/137,036

22 <151> PRIOR FILING DATE: 2002-04-30

24 <150> PRIOR APPLICATION NUMBER: U.S. No. 09/724,624

25 <151> PRIOR FILING DATE: 2000-11-28

27 <150> PRIOR APPLICATION NUMBER: U.S. No. 09/598,401

28 <151> PRIOR FILING DATE: 2000-06-20

30 <150> PRIOR APPLICATION NUMBER: PCT/NZ00/00018

31 <151> PRIOR FILING DATE: 2000-02-24

33 <150> PRIOR APPLICATION NUMBER: U.S. No. 60/146,591

34 <151> PRIOR FILING DATE: 1999-07-30

36 <150> PRIOR APPLICATION NUMBER: U.S. Patent No. 09/276,599

37 <151> PRIOR FILING DATE: 1999-03-257
39 <160> NUMBER OF SEQ ID NOS 125) Include intertorelly Skipped 5 equires
41 <170> SOFTWARE: FastSEQ for Windows Version 4.0 in (1607 response

ERRORED SEQUENCES

E--> 2927 <210> SEQ ID NO: 125

(see p.2)

<400> 120					
aaaaaaggga aacattatac	caaattttat	datatettee	aacaacatac	tettetatat	60
atggtgcctc ctctgatgga					120
					180
gcccccactc acaagataat					
cttttgatgt accgagtaca	ctactcatgg	tgaaggccgt	gtcttgcagc	ttttcccatg	240
gtttattttg aaagtaatag	tactggacct	catttgcaac	gacacataat	attcttactg	300
acgacacttt gtttgatttc	ttatagaaaa	atgcaaggtg	gcacaaaaag	atggaaagcc	360
cgacctatca agcatacgaa	gggtcatgtt	cacaccctct	gaaatcttca	gagtctcacc	420
ctatgttgga cgctaatcaa					480
aatcaatcat tgaaaaatat	accagataac	tcctacgatg	gaggggatta	tttgcgtacc	540
ctccgcgtgg gtgggcacat	tgggcaggtc	ctttggtaag	tcttggagac	agagtcacgt	600
ttccataatt gaagtggaca	tttatgaatc	tttcgaaagt	tgtagaactc	ttaattttcg	660
acggaatagt ttgacacgtt	ttgtacgatc	tggtttttcc	ggggaacgcc	aattttggtt	720
tctgaaggac agcatttaca	atattgtctg	tcgttgacca	ggacagctgg	ctcggaactc	780
gggtttccga tgcgcaggaa	gcgcattgaa	atgagaatat	aatctagttc	tacctgtgga	840
gctatcacaa aatactaaaa	ctggtggaca	tacctcttgt	ctgttctcga	aatcggccaa	900
aatgggaaag aagagggtag	agctgaaacg	cattcaaaac	cctagcagtc	gacatgctac	960
tttctctaaa cgcaagaatg					1020
tgctgaagtc gctctcatca					1080
taacgatatg gcagcaattc					1140
cagtccaaca tcgcttcaaa	acgtaaagta	tcatgaatca	gggcttgaga	aattgcaaga	1200
gaagttgacc gctttgcaaa					1260
aacaatgaaa gaactgcaac					1320
					1326
Jequeres 121	though	124 m	user		

<210> 125

see P.3

3

RAW SEQUENCE LISTING ERROR SUMMARY

PATENT APPLICATION: US/10/702,319

DATE: 11/17/2003 TIME: 09:26:38

Input Set : A:\1036C5SEQLIST.txt

Output Set: N:\CRF4\11172003\J702319.raw

Skipped Sequences (NEW RULES):

Sequence(s) missing. If intentional, please use the following format for each skipped sequence.

<210> sequence id number <400> sequence id number 000

Seq#:121,122,123,124

sel p.4 for more enon

10/702,319

<210> 3 <211> 1226 <212> DNA <213> Pinus radiata

<221> 5'UTR <222> (1)...(1266) (1226) orby 1226 huelevirdir in the seguence

VERIFICATION SUMMARY

DATE: 11/17/2003 TIME: 09:26:38 PATENT APPLICATION: US/10/702,319

Input Set : A:\1036C5SEQLIST.txt

Output Set: N:\CRF4\11172003\J702319.raw

L:15 M:270 C: Current Application Number differs, Replaced Current Application No L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:55 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1 L:58 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1 L:61 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ 1D#:1 L:175 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2 L:219 M:351 W: Sequence data Name/Key Feature Out-of-Range, SEQ ID#:3, 5UTR LOCATION: (1)... (1266)L:256 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4 L:259 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4 L:282 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5 L:301 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6 L:304 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6 L:328 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ 1D#:7 L:544 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:480 L:2927 M:216 E: (34) Seq. #s missing, SEQ 1D NOS: 121 thru 124